

Steven M. Ruben  
Appl. No. 10/662,429

Department \_\_\_\_\_

Subject \_\_\_\_\_

Name GUO-Liang YU

Address \_\_\_\_\_

National® Brand

# Computation Notebook

11<sup>3</sup>/<sub>4</sub>" x 9<sup>1</sup>/<sub>4</sub>", 4 x 4 Quad., 75 Sheets

43-648



0 73333 43648 8



AVERY  
DENNISON

Office Products  
Chicopee, MA 01022

Ruben EXHIBIT #63

Department \_\_\_\_\_

Subject \_\_\_\_\_

Name GUO-Liang YU

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# Computation Notebook

11 3/4" x 9 1/4", 4 x 4 Quad., 75 Sheets

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DENNISON

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Ruben EXHIBIT 2063  
Ruben v. Wiley et al.  
Interference No. 105,077  
RX 2063

1/23/96 Large scale plasmid prep for cDNA library

HBC to be used to make single stranded DNA

to test SAGE

— transformation from an old stock of plasmid DNA

— obtain titer  $1.5 \times 10^7$  cfu grow up to 1 L O/W

— plasmid prep

spin down, resuspend in 40 ml H<sub>2</sub>O  
add 8 ml NaOH / SDS  
60 ml 3M KOAC

spin down ppt with isopropanol

CsCl<sub>2</sub> gradient 60K g/h

Concentration of DNA is low

2

# 1/24 proliferation assay for endothelial cells

## EDAP

cell lines Arteray endothelial cell  
Veinus endothelial cells  
Huvec

Seed  $5 \times 10^3$  cells to each 96 well DMEM 10% FBS  
5 hr incubation — let cell attach, remove medium  
add Samples as show below

	TNF $\alpha$				EDAP E2				EDAP E3			
	1	2	3	4	5	6	7	8	9	10	11	12
A		1 ng			1 ng			1 ng			10%	✓
B		100 ng			100 ng			100 ng			1%	✓
C		10 ng			10 ng			10 ng			0.1%	✓
D		1 ng			1 ng			1 ng			0.01%	✓
E		bfr			1 ng			100 ng			10%	✓
F					100 ng			10 ng			1%	✓
G		Control			10 ng			1 ng			0.1%	✓
H					1 ng			0.1 ng			0.01%	✓

EDAP  
E4

bfr

40 hr after TNF $\alpha$  assay for AUC & VE  
add  $^3\text{H}$  thymidine  
incubate 6 hrs  
wash once w/ PBS. Lyse cell Count

# Certificate of Analysis

3

Protein Name: TNF gamma (EDAP)

Batch #: HG02704-E2

Date: 11/10/95

Expression System: E. Coli

## Method of Purification:

1. Cell disruption and purification of inclusion body  
Wash pellet with TE, 1M, 2 M Urea, and
2. Extraction TNF  $\gamma$  with 4 and 8 M Urea
3. Buffer exchange (2XPBS) and refolding in PD-10
4. Removing endotoxin by affinity gel chromatography

Estimated Purity: > 80%



TNF gamma

Protein Concentration:

Method A: 176  $\mu$ g/ml

Method B: 180  $\mu$ g/ml

Endotoxin level in sample: 0.17 EU/mg

Method of Analysis: Limulus Amebocyte Lysate test (Bio-Whittaker)

Sterile filtered: Yes ☒ No ☐

Buffer composition: 2XPBS

Total amount of protein purified: 14 mg

Prepared by: Jian Ni

Approved by: *Kevin Jeff*

*1.5nd 5/3 11/30*

Updated May 23, 1995

# Certificate of Analysis

Protein Name: TNF gamma

Batch #: HG02703-3E

*43 5/5 ~ 5/6*

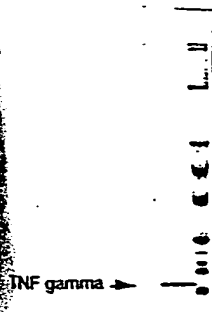
Date: 5/26

Expression System: E. Coli

## Method of Purification:

1. Cell disruption and purification of inclusion body
2. Solubilization of protein with 8 M urea
3. Sephadex S-200 size exclusion chromatography
4. Dialysis into PBS

Estimated Purity: > 70%



TNF gamma

Protein Concentration: 80  $\mu$ g/ml

Endotoxin level in sample: 525 EU/mg

Method of Analysis: Limulus Amebocyte Lysate test (Bio-Whittaker)

Buffer composition: PBS

Endotoxin level in buffer: <10 EU/ml

Sterile filtered: Yes ☒ No ☐

Prepared by: Jian Ni

Approved by: *Kevin Jeff*

Updated May 23, 1995

## Certificate of Analysis

Protein Name: TNF-delta (His-tag)  
Clone ID#: HLTBT71

Batch #: HG10700-E1

Date: 1/22/96

Expression System: E. Coli

Molecular weight: 20.8 Kd

Method of Purification:  
Nickel-chelate affinity chromatography column  
Buffer exchange (2XPBS) and refolding in PD-10 column

Estimated Purity: 80%



Protein Concentration:

Method A: 200 µg/ml

Method B: 160 µg/ml

Endotoxin level in sample: 2.8 EU/mg

Method of Analysis: Limulus Amebocyte Lysate test (Bio-Whittaker)

Sterile filtered: Yes Y No   

Buffer composition: 2XPBS

Biological activity  
yes/not tested

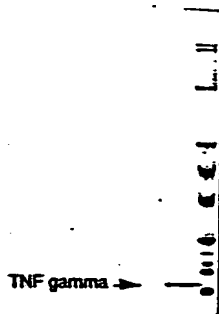
Total amount of protein purified: 10 mg

Prepared by: Jian Ni

Approved by:

Updated September 15, 1995  
by replacement of 20.0 size exclusion chromatography  
4, Dialysis into PBS

Estimated Purity: > 70%



Protein Concentration: 80 µg/ml

Endotoxin level in sample: 525 EU/mg

Method of Analysis: Limulus Amebocyte Lysate test (Bio-Whittaker)

Buffer composition: PBS

Endotoxin level in buffer: <10 EU/ml

Sterile filtered: Yes XXX No   

Prepared by: Jian Ni

Approved by:

*Jian Ni*

Updated May 21, 1995

## Certificate of Analysis

3

In Name: TNF gamma (EDAP)

#: HG02704-E2

Date: 11/10/95

ssion System: E. Coli

d of Purification:  
disruption and purification of inclusion body  
elut with TE, 1M, 2 M Urea, and  
ction TNF γ with 4 and 8 M Urea  
r exchange (2XPBS) and refolding in PD-10  
oving endotoxin by affinity gel chromatography

ted Purity: > 80%



TNF gamma

Concentration:

hod-A: 176 µg/ml

hod B: 180 µg/ml

level in sample: 0.17 EU/mg

Analysis: Limulus Amebocyte Lysate test (Bio-Whittaker)

## Certificate of Analysis

Protein Name: TNF-gamma (EDAP) (His-tag)  
Clone ID#: HUVEO91

Batch #: HG02702-E4

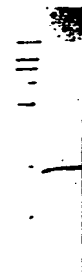
Date: 1/22/96

Expression System: E. Coli

Molecular weight: 20.1 Kd

Method of Purification:  
Nickel-chelate affinity chromatography column  
Buffer exchange (2XPBS) and refolding in PD-10 column

Estimated Purity: >80%



Protein Concentration:

Method A: 150 µg/ml

Method B: 260 µg/ml

Endotoxin level in sample: 3.1 EU/mg

Method of Analysis: Limulus Amebocyte Lysate test (Bio-Whittaker)

Sterile filtered: Yes Y No   

Buffer composition: 2XPBS

Biological activity  
yes/not tested

Total amount of protein purified: 10 mg

Prepared by: Jian Ni

Approved by:

Updated September 15, 1995

4

MTS assay:

AE

	1	2	3	4	5	6	7	8	9	10	11	12
A	0.019	0.033	0.070	0.033	0.026	0.013	0.006	0	0.012	0.011	0.011	
B	0.47	0.043	0.071	0.058	0.050	0.043	0.047	0.042	0.048	0.162	0.162	
C	0.039	0.01	0.073	0.040	0	0	0.046	0.04	-0.005	0.018	0.018	
D	-0.051	0.011	0.070	0.014	0	0	0	-0.04	-0.019	0	0	
E	-0.027	0.038	0.043	0.010	-0.029	0.017	0.082	0.077	0.070	0.289	0.289	
F	-0.02	0.066	0.045	0.084	0.033	0.04	0.145	0.133	0.09	0.212	0.212	
G	0.015	0.04	0.028	0.02	0.007	0	0.086	0.091	0.1	0.059	0.059	
H	-0.01	0.061	0.072	0.09	0.039	0.001	0.042	0.013	0.041	0.017	0.017	

b5

E4

BFGF

FBS

	1	2	3	4	5	6	7	8	9	10	11	12
Limit												
A Absorb	-0.005	-0.013	-0.008	0.007	0.039	0.050	0.021	0.017	0.001	-0.004	0.658	-0.020
Range												
Limit												
B Absorb	-0.034	-0.040	-0.008	0.003	0.008	0.030	0.024	0.021	0.019	0.023	0.658	-0.013
Range												
Limit												
C Absorb	-0.028	-0.027	-0.007	-0.010	0.027	0.027	0.024	0.035	0.096	0.040	0.121	-0.008
Range												
Limit												
D Absorb	-0.025	-0.003	0.010	0.015	0.023	0.028	0.026	0.030	0.024	0.034	0.050	-0.002
Range												
Limit												
E Absorb	-0.024	-0.014	-0.005	-0.016	0.013	0.014	0.016	0.309	0.237	0.269	0.847	0.000
Range												
Limit												
F Absorb	-0.074	-0.060	-0.035	-0.032	-0.030	-0.008	0.011	0.302	0.331	0.395	0.646	-0.013
Range												
Limit												
G Absorb	-0.042	-0.022	-0.014	-0.018	0.002	-0.013	0.009	0.171	0.191	0.213	0.158	-0.006
Range												
Limit												
H Absorb	-0.027	-0.025	-0.018	0.005	0.018	0.019	0.004	0.090	0.094	0.069	0.042	0.000
Range												

VE

Huvec

AE

USER: 3 ID:34 CPM PRESET TIME: 1.00  
SAMPLE REPEAT: 1 CYCLE REPEAT: 1 SCR:M RS232:N  
NR: 1 AOC:N QCF:N RCM:Y  
RCH-TIME: 0.50 INT:999.95  
CHANNEL 1-LL: 0 UL: 400 ZSIGMA: 0.50 BKG SUB: 0.00 BKG ZSIG  
DATA CALC: CPM, UNKNOWN REPLICATES: 1 NORM FACTOR:0 1.000000  
HALF LIFE(DAYS):N

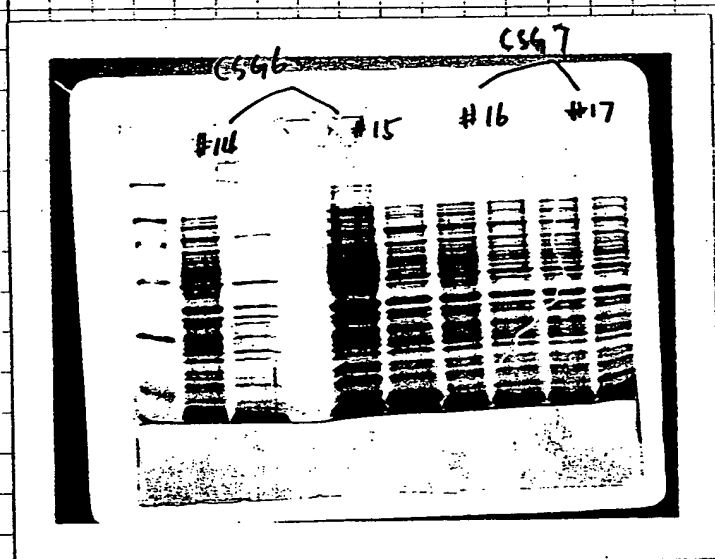
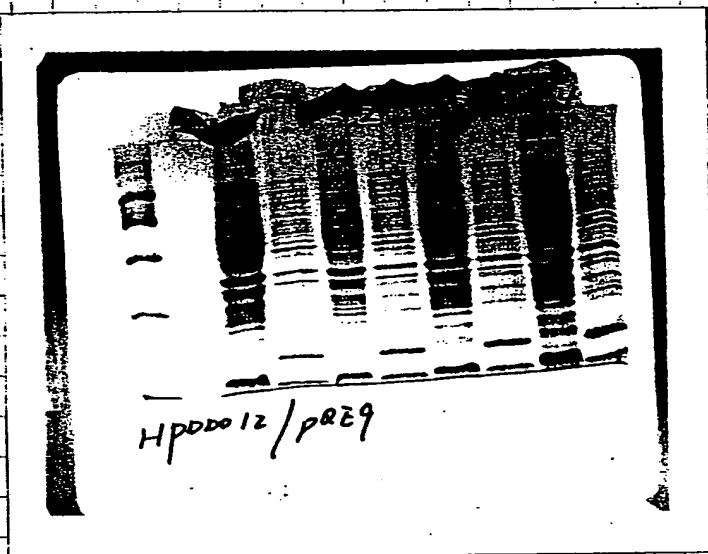
TUE 15 M

SAM	POS	CH	CPM	ZSIGZ	TIME	EL	TIME	AVG	H#	RCMZ										
1	163	1	253.00	12.57	1.00	1.19	62.0			18.31	86 248	1	37.00	21.44	1.00	107.04	59.0	19.57		
2	164	1	1552.00	5.08	1.00	2.42	61.0			1.30	87 249	1	175.00	15.12	1.00	108.28	55.0	53.38		
3	165	1	141.00	16.84	1.00	3.67	64.0			21.97	88 250	1	76.00	22.94	1.00	109.52	53.0	33.55		
4	166	1	301.00	11.33	1.00	4.90	66.0			1.30	89 251	1	59.00	26.04	1.00	110.75	53.0	39.26		
5	167	1	397.00	10.04	1.00	6.14	64.0			21.97	90 252	1	70.00	23.90	1.00	111.98	52.0	39.88		
6	168	1	223.00	13.39	1.00	7.38	64.0			18.88	91 253	1	50.00	28.28	1.00	113.22	52.0	44.14		
7	169	1	363.00	10.47	1.00	8.61	63.0			10.93	92 254	1	41.00	31.23	1.00	114.45	51.0	23.34		
8	170	1	336.00	10.91	1.00	9.85	64.0			21.35	93 255	1	44.00	30.15	1.00	115.68	52.0	39.68		
9	171	1	186.00	14.66	1.00	11.10	60.0			11.71	94 256	1	147.00	16.30	1.00	116.92	52.0	33.02		
10	172	1	116.00	18.57	1.00	12.34	63.0			23.55	95 257	1	56.00	26.73	1.00	118.17	50.0	26.20		
11	173	1	266.00	12.26	1.00	13.57	62.0			12.63	96 258	1	52.00	27.74	1.00	119.40	50.0	51.11		
12	174	1	633.00	7.94	1.00	14.80	64.0			14.96	97 259	1	108.00	19.25	1.00	120.64	53.0	29.10		
13	175	1	98.00	20.20	1.00	16.04	63.0			3.46	98 260	1	163.00	15.67	1.00	121.87	52.0	29.10		
14	176	1	233.00	13.10	1.00	17.28	70.0			18.38	99 261	1	55.00	27.47	1.00	123.11	51.0	29.20		
15	177	1	164.00	15.62	1.00	18.52	63.0			32.91	100 262	1	35.00	27.47	1.00	124.34	52.0	23.08		
16	178	1	36.00	26.73	1.00	19.74	57.0			16.88	101 263	1	55.00	26.97	1.00	125.59	59.0	50.81		
17	179	1	112.00	18.90	1.00	20.97	60.0			20.00	102 264	1	138.00	17.03	1.00	126.82	52.0	30.66		
18	180	1	101.00	19.90	1.00	22.21	59.0			14.87	103 265	1	34.00	34.30	1.00	128.05	52.0	35.53		
19	181	1	1932.00	4.35	1.00	23.44	59.0			15.98	104 266	1	42.00	30.86	1.00	129.29	53.0	38.72		
20	182	1	962.00	6.45	1.00	24.67	61.0			1.18	105 267	1	66.00	24.62	1.00	130.54	58.0	63.75		
21	183	1	357.00	10.59	1.00	25.92	64.0			3.14	106 268	1	81.00	22.22	1.00	131.78	53.0	33.25		
22	184	1	195.00	14.32	1.00	27.16	67.0			27.14	107 269	1	119.00	18.33	1.00	133.02	57.0	50.75		
23	185	1	687.00	7.63	1.00	28.40	67.0			38.97	108 270	1	92.00	20.85	1.00	134.25	51.0	46.79		
24	186	1	334.00	10.63	1.00	29.65	63.0			8.14	109 271	1	168.00	15.43	1.00	135.50	59.0	52.65		
25	187	1	177.00	13.03	1.00	30.88	65.0			14.77	110 272	1	130.00	17.54	1.00	136.75	58.0	50.21		
26	188	1	261.00	12.38	1.00	32.13	66.0			12.05	111 273	1	188.00	14.99	1.00	138.00	59.0	61.70		
27	189	1	248.00	12.70	1.00	33.36	69.0			19.46	112 274	1	45.00	29.81	1.00	139.24	51.0	34.91		
28	190	1	238.00	12.96	1.00	34.59	64.0			5.22	113 275	1	61.00	25.61	1.00	140.47	54.0	39.45		
29	191	1	477.00	9.16	1.00	35.81	66.0			16.67	114 276	1	45.00	29.81	1.00	141.70	52.0	26.28		
30	192	1	325.00	11.09	1.00	37.08	67.0			25.74	115 277	1	134.00	16.12	1.00	142.95	60.0	61.37		
31	193	1	337.00	10.59	1.00	38.31	66.0			8.05	116 278	1	54.00	27.22	1.00	144.18	50.0	40.62		
32	194	1	366.00	10.45	1.00	39.56	66.0			27.29	117 279	1	29.00	37.14	1.00	145.41	52.0	28.05		
33	195	1	351.00	10.68	1.00	40.80	65.0			22.51	118 280	1	99.00	20.10	1.00	146.65	56.0	42.92		
34	196	1	242.00	12.66	1.00	42.03	65.0			7.28	119 281	1	37.00	32.88	1.00	147.88	60.0	27.07		
35	197	1	325.00	11.09	1.00	43.28	65.0			31.25	120 282	1	51.00	28.01	1.00	149.12	61.0	27.62		
36	198	1	436.00	9.56	1.00	44.53	64.0			24.16	121 283	1	63.00	25.20	1.00	150.35	57.0	23.21		
37	199	1	473.00	9.20	1.00	45.77	70.0			6.82	122 284	1	60.00	25.82	1.00	151.58	61.0	31.22		
38	200	1	428.00	9.67	1.00	47.00	66.0			3.95	123 285	1	67.00	24.43	1.00	152.82	53.0	24.78		
39	201	1	297.00	11.61	1.00	48.23	69.0			10.21	124 286	1	65.00	24.81	1.00	154.05	56.0	29.27		
40	202	1	318.00	11.22	1.00	49.48	68.0			30.33	125 287	1	47.00	29.17	1.00	155.27	57.0	16.35		
41	203	1	348.00	10.72	1.00	50.73	70.0			23.16	126 288	1	45.00	29.81	1.00	156.51	61.0	18.46		
42	204	1	231.00	13.16	1.00	51.97	65.0			35.15	127 289	1	59.00	26.04	1.00	157.73	56.0	13.03		
43	205	1	394.00	10.08	1.00	53.22	67.0			25.10	128 290	1	81.00	22.22	1.00	158.97	56.0	48.20		
44	206	1	892.00	6.79	1.00	54.47	70.0			13.91	129 291	1	104.00	19.61	1.00	160.21	58.0	34.80		
45	207	1	115.00	18.65	1.00	55.71	67.0			61.78	130 292	1	77.00	22.79	1.00	161.45	53.0	42.13		
46	208	1	1239.00	5.68	1.00	56.96	68.0			6.30	131 293	1	144.00	16.67	1.00	162.68	53.0	13.50		
47	209	1	553.00	8.50	1.00	58.20	67.0			132 294	1	60.00	25.82	1.00	163.91	56.0	33.86			
48	210	1	244.00	12.80	1.00	59.45	71.0			133 295	1	53.00	27.47	1.00	165.15	61.0	32.19			
49	211	1	266.00	12.26	1.00	60.69	67.0			23.42	134 296	1	60.00	25.82	1.00	166.38	56.0	32.51		
50	212	1	433.00	9.59	1.00	61.92	69.0			15.31	135 297	1	115.00	18.65	1.00	167.62	59.0	60.08		
51	213	1	289.00	11.76	1.00	63.16	64.0			19.32	136 298	1	102.00	19.80	1.00	168.86	61.0	58.56		
52	214	1	197.00	14.25	1.00	64.40	69.0			137 299	1	40.00	31.62	1.00	170.10	61.0	14.29			
53	215	1	243.00	12.83	1.00	65.64	67.0			138 300	1	43.00	30.50	1.00	171.33	59.0	20.39			
54	216	1	333.00	10.96	1.00	66.89	67.0			139 301	2	34.00	34.30	1.00	172.56	62.0	28.12			
55	217	1	143.00	16.72	1.00	68.13	68.0			140 302	3	82.00	22.09	1.00	173.79	56.0	31.15			
56	218	1	183.00	14.78	1.00	69.37	67.0			141 303	4	50.00	28.28	1.00	175.03	54.0	39.19			
57	219	1	6383.00	2.47	1.00	70.61	67.0			142 304	5	72.00	23.57	1.00	176.27	55.0	57.92			
58	220	1	907.00	6.64	1.00	71.85	64.0			143 305	6	69.00	24.08	1.00	177.49	54.0	23.80			
59	221	1	470.00	9.23	1.00	73.09	64.0			144 306	7	93.00	20.74	1.00	178.73	60.0	41.00			
60	222	1	333.00	10.93	1.00	74.33	64.0			145 307	8	47.00	29.17	1.00	179.97	53.0	31.18			
61	223	1	303.00	11.49	1.00	75.57	64.0													





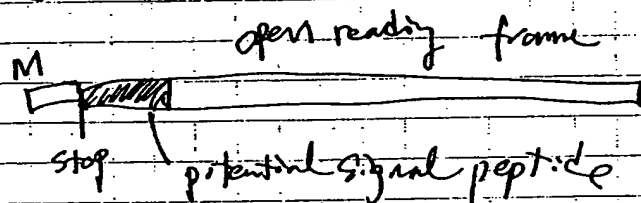
TNF $\alpha$  expression in pOE9 His tagged



3/12/96 TNF Receptor like HTBN61

Now called Death Domain containing Receptor DDCR

Full length clone HLMH058 contained a stop codon after methionine



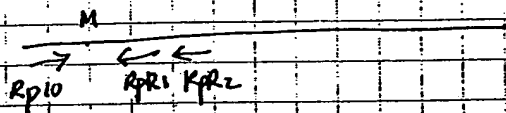
the stop codon is real: with many rounds of sequencing

① to get new clones

② PCR 5' seq direct sequence

oligo HTBN61Rp10 made

CCCTCCGAAACCTGGTG



use Rp10 and RpR2 to PCR libraries. Sequence the product with RpR1

3/13/98 Expression vectors:

① EDAP  $\Delta 3$   $\delta 48$  5' CGCCATGGCCTTCACCAAGAACCGA NcoI

② EDAP  $\Delta 3$   $\delta 54$  5' CGCCCATGGACTATACCAASAAATTC NcoI

both clone into pDE60 3' will use EDAP HindIII

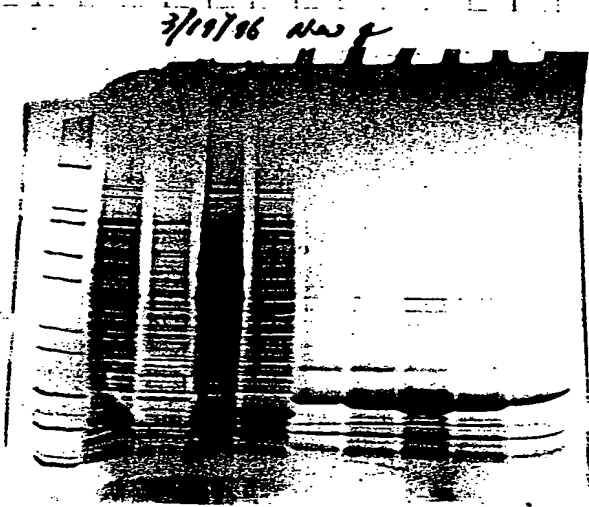
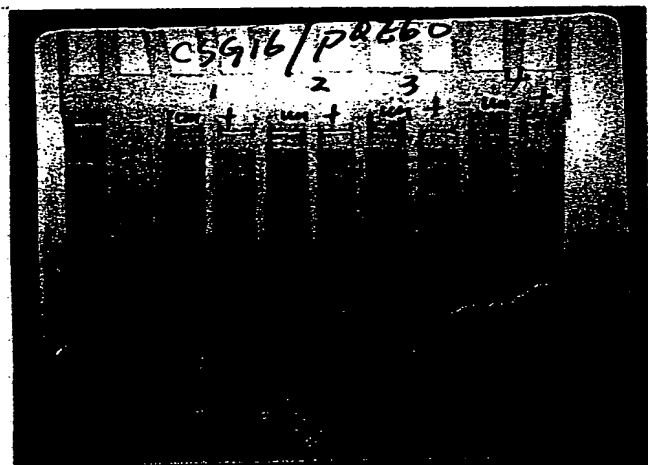
③ AIM2 $\Delta 1$  5' CGCCATGGCCAACTCCAGCTTGACC NcoI

④ AIM2 $\Delta 2$  5' CGCCATGGTCA~~CC~~CGCCTGCTG~~CA~~ NcoI

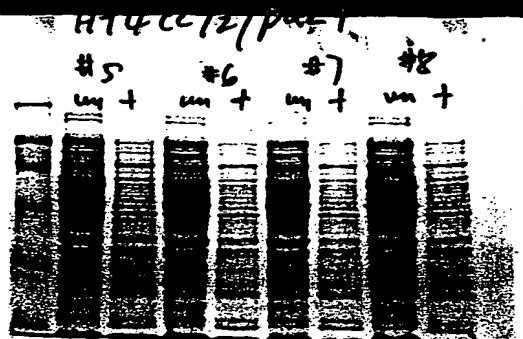
both clone into pDE60 3' HindIII

SUPERVISOR  
DATE

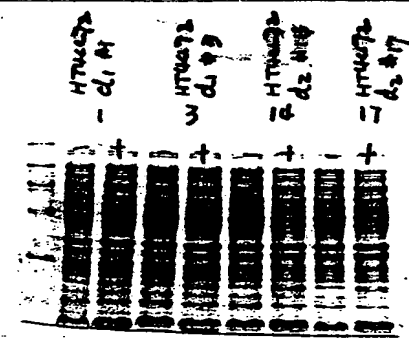
SUPERVISOR  
DATE



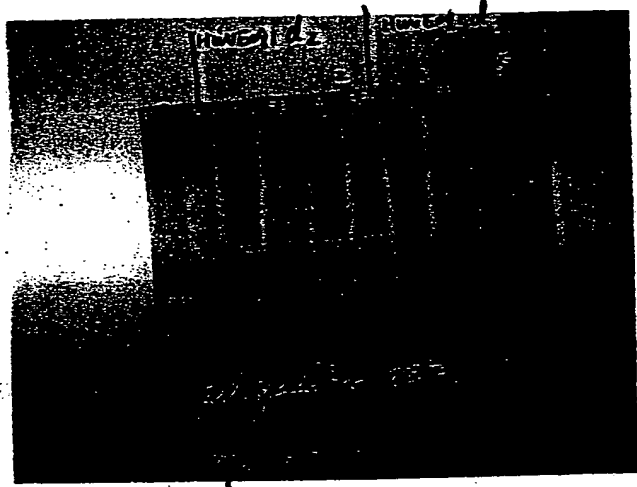
Anti-His CS8



3/15/96



all gene in PRE60  
10% SOS



EDAP Expression vector position

781 ATATGAGACGCTTTTAAAGCAAAGTCTACAGTTTCCCAATGAGAAAATTAATCCTCTTTC  
M R R F L S K V Y S F P M R K L I L F L  
PQE9, PQE70, CHO  
841 TTGTCTTTCCAGTTGTGAGACAAACTCCACACAGCACTTTAAAAATCAGTTCCAGCTC  
V F E V Y R Q T P T Q H F K N Q F P A L  
IL6 PQE60, pA2-GP  
901 TGCACGTGGGAACATGAAGTGGCCTTCCACCAAGAACCGAATGACTATACCAACA  
H W E H E L G L A F T K N R M N Y T N K  
PQE60  
961 AATTCCTGCTGATCCAGAGTCGGGAGACTACTTCATTTACTCCAGGTCACATTCCGTG  
F L L I P E S G D Y F I Y S Q V T F R G  
1021 GGATGACCTCTGAGTGCAGTGAAATCAGACAAGCAGCCGACCAACAAGCCAGACTCCA  
M T S E C S E I R Q A G R P N K P D S I  
1081 TCACTGTGGTCATCACCAGGTAAACAGACAGCTACCCTGAGCCAACCCAGCTCCTCATGG  
T V V I T K V T D S Y P E P T Q L L M G  
1141 GGACCAAGTCTGTATGCGAAGTAGGTAGCAACTGGTTCCAGCCCATCTACCTCGGAGCCA  
T K S V C E V G S N W F Q P I Y L G A M  
1201 TGTTCCTTGAAGAAGGGGACAAGCTAATGGTGAACGTCAGTGACATCTCTTTGGTGG  
F S L Q E G D K L M V N V S D I S L V D  
1261 ATTACACAAAAGAAGATAAAACCTTCTTTGGAGCCTTCTTACTATASGAGGAGAGCAAAT  
Y T K E D K T F F G A F L L \*

538 d1  
448 d2  
454 d3

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